

ABSTRACT

Novel enterokinase cleavage sequences are provided. Also disclosed are methods for the rapid isolation of a protein of interest present in a fusion protein construct including a novel enterokinase cleavage sequence of the present invention and a ligand recognition sequence for capturing the fusion construct on a solid substrate. Preferred embodiments of the present invention show rates of cleavage up to thirty times that of the known enterokinase cleavage substrate (Asp)₄-Lys-Ile.

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